

BLAST PubMed Nucleotide Protein Genome Structure PopSet Taxonomy Help

FASTA view

Sequence feature view of the region: (gi|36791:829-924, 970-1074, 1120-1209, 1255-1374, 1420-1581, 1691-1798, 1843-1890, 2054-2134)

```
LOCUS
            36791
                           810 bp
                                                      PRI
                                                                30-S
DEFINITION
            CDS from: Homo sapiens TCF-1 gene.
ACCESSION
            36791
VERSION
            36791
KEYWORDS
            T-cell transcription factor; transcription factor.
SOURCE
            human.
  ORGANISM
            Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eu
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; H
REFERENCE
               (bases 829 to 2134)
  AUTHORS
            van de Wetering, M., Oosterwegel, M., Holstege, F., Dooye
            Suijkerbuijk, R., Geurts van Kessel, A. and Clevers, H.
  TITLE
            The human T cell transcription factor-1 gene. Structur
            localization, and promoter characterization
  JOURNAL
            J. Biol. Chem. 267 (12), 8530-8536 (1992)
  MEDLINE
            92235082
REFERENCE
            2
                (bases 829 to 2134)
  AUTHORS
            Van de Wetering, M., Castrop, J., Korinek, V. and Clevers
  TITLE
            Extensive alternative splicing and dual promoter usage
            Tcf-1 protein isoforms with differential transcription
            properties
  JOURNAL
            Mol. Cell. Biol. 16 (3), 745-752 (1996)
  MEDLINE
            96182076
REFERENCE
               (bases 829 to 2134)
  AUTHORS
            Van de Wetering, M.L.
  TITLE
            Direct Submission
  JOURNAL
            Submitted (07-JAN-1992) M.L. Van De Wetering, Departme
            Immunology, P.O. Box 8550, 3508 GA Utrecht, THE NETHER
FEATURES
                     Location/Qualifiers
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                         276 c
                                  200 g
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      121 ggtgtcccc aactctctct ctacgaacat ttcaacagcc cacatcccac c
      181 gcggacatca gccagaagca agttcacagg cctctgcaga cccctgacct c
      241 tactccctga cctcaggcag catggggcag ctccccaca ctgtgagctg g
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      481 ttcatqctqt acatgaagga gatgagagcc aaggtcattg cagagtgcac a
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      601 gccaagtact atgagctggc ccgcaaggag aggcagctgc acatgcagct a
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```